

M	C	P	S	Q	P	T	Y	P	G	D	P	G	P
ATG	TGC	CCG	AGC	CAG	CCG	ACC	TAT	CCG	GGC	GAT	CCC	GGG	CCG
V	E	D	L	I	R	F	Y	D	N	L	Q	Q	W
GTG	GAA	GAT	CTG	ATC	CGC	TTT	TAT	GAT	AAC	CTG	CAG	CAG	TGG
L	N	C	V	T	A	A	C	*					
CTG	AAC	TGC	GTG	ACC	GCC	GCC	TGC	TAG					

FIGURE 1

10027038-122004

1	11	21	31	41
ACACACCATA	TGTGCCCAG	CCAGCCGACC	TATCCGGGCG	ATCCCCGGCC
TGTGTGGTAT	ACACGGGCTC	GGTCGGCTGG	ATAGGCCCGC	TAGGGCCCCG
51	61	71	81	91
GGTGAAGAT	CTGATCCGCT	TTTATGATAA	CCTGCAGCAG	TGGCTGAACT
CCACCTTCTA	GACTAGGCGA	AAATACTATT	GGACGTCGTC	ACCGACTTGA
101	111	121	131	
GGGTGACCGC	CGCCTGCTAG	GGATCCACAC	AC	
CGCACTGGCG	GCGGACGATC	CCTAGGTGTG	TG	

FIGURE 2

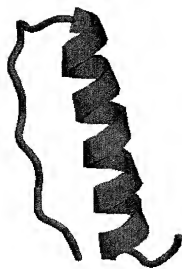


FIGURE 3

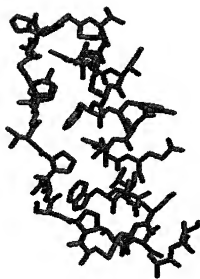


FIGURE 4



FIGURE 5

10027038.122001

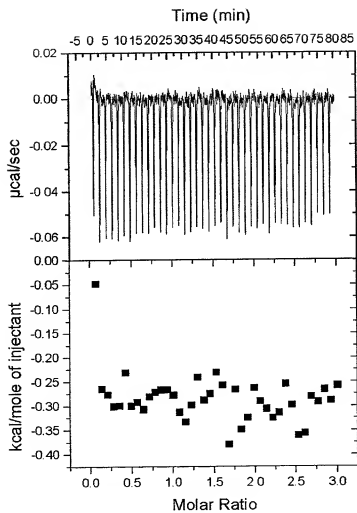


FIGURE 6

10027038.122001
10027038.122001

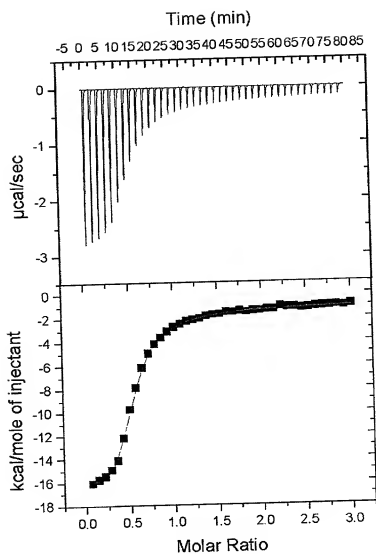


FIGURE 7

10027036 122001

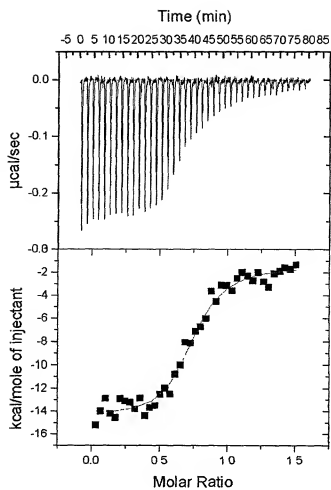


FIGURE 8

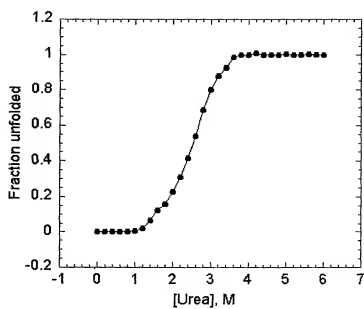


FIGURE 9

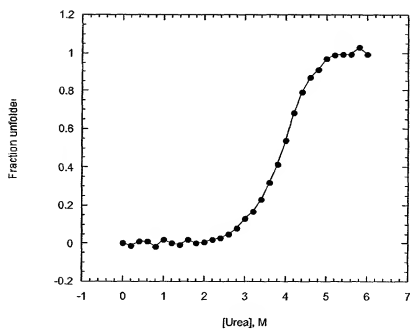


FIGURE 10

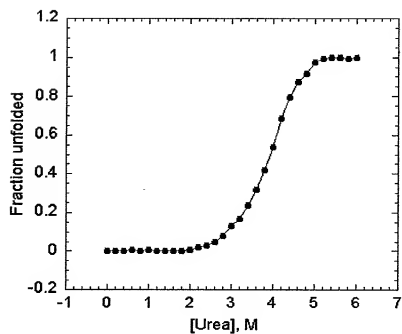


FIGURE 11

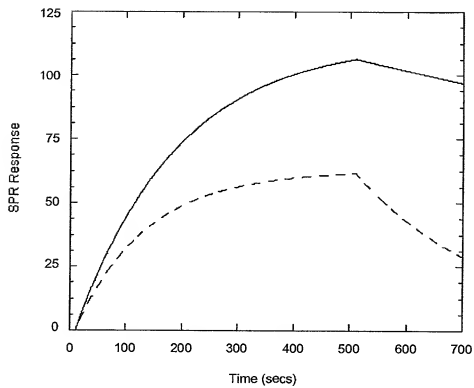


FIGURE 12

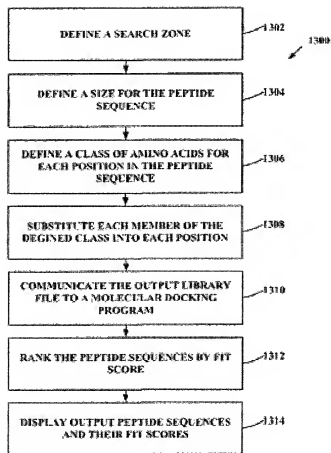


FIG. 13

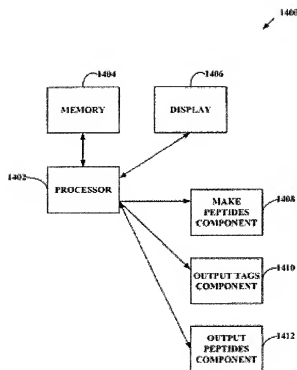


FIG. 14

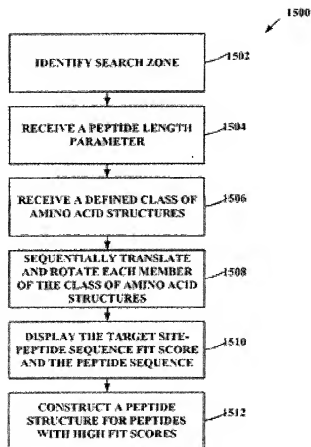


FIG. 15